

32. The enzyme of claim ~~27~~<sup>1</sup>, which has an amino acid sequence that is at least 98% identical to amino acids 26-485 or 26-646 of SEQ ID NO: 10.
33. The enzyme of claim ~~27~~<sup>1</sup>, which comprises an amino acid sequence of amino acids 26-485 of SEQ ID NO: 10.
34. The enzyme of claim ~~27~~<sup>1</sup>, which comprises an amino acid sequence of amino acids 26-646 of SEQ ID NO: 10.
35. The enzyme of claim ~~27~~<sup>1</sup>, which is encoded by the endoglucanase encoding part of the DNA sequence obtainable from the plasmid in *Escherichia coli* DSM 12805.
36. The enzyme of claim ~~27~~<sup>1</sup>, which is a *Bacillus licheniformis* enzyme.
37. The enzyme of claim ~~26~~<sup>14</sup>, which is a *Bacillus licheniformis*, ATCC 14580 enzyme.
38. The enzyme of claim ~~27~~<sup>1</sup>, which is active at a pH in the range of 4-11.
39. The enzyme of claim ~~26~~<sup>12</sup>, which is active at a pH in the range of 5.5-10.5.
40. An enzyme composition comprising the enzyme of claim ~~27~~<sup>1</sup>.
41. The composition of claim ~~40~~<sup>14</sup>, which further comprises one or more enzymes selected from the group consisting of alpha-amylases, cellobiohydrolases, cellulases (endoglucanases), cutinases, beta-glucanases, glucoamylases, hemicellulases, laccases, ligninases, lipases, oxidases, pectate lyases, pectin acetyl esterases, pectinases, pectin lyases, pectin methylesterases, peroxidases, phenoloxidases, polygalacturonases, proteases, pullulanases, reductases, rhamnogalacturonases, xylanases, xyloglucanases, other mannanases, transglutaminases; and mixtures thereof.
42. A method for degradation of cellulose-containing biomass, comprising treating the biomass with an effective amount of the enzyme of claim ~~27~~<sup>1</sup>.